

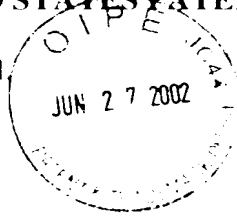
**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE**

Application of: Kessler, et al.

Serial No.: 09/530,746

Filed: November 16, 2000

For: **SPECIFIC AND SENSITIVE METHOD  
FOR DETECTING NUCLEIC ACIDS**



Group Art Unit: 1654

Examiner: Strzelecka, T.

Attorney Docket No.: 1803-277-999

**AMENDMENT UNDER 37 C.F.R. § 1.111**

Commissioner for Patents  
Washington, D.C. 20231

Sir:

In response to the Notice to Comply with the Sequence Rules mailed September 7, 2001, please amend the above-identified application. This amendment is accompanied by (1) Appendix A showing a marked-up version of the amended claim; (2) Appendix B showing a clean copy of the claims as pending after entry of this amendment; (3) a Petition Pursuant to 37 C.F.R. § 1.137(b) to revive an unintentionally abandoned patent application; and (4) a Sequence Listing in compliance with 37 C.F.R. §§ 1.821-1.825 in paper and computer readable format.

**AMENDMENT**

**IN THE CLAIMS:**

Please amend the claim to read as follows:

22. (Twice Amended) A method for the detection of HCV, wherein two primers and one probe are used whose sequences are derived from sequences of consecutive bases the HGBV sequence shown below, complementary sequences thereto or sequences that are more than 80% identical to these sequences:

5'-GTACTGCCTGATAGGGTCCTTGCGAGGGGATCTGGGAGTC  
TCGTAGACCG TAGCACATG-3' (SEQ ID NO: 95).

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